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Gly Thr Trp Gly Ile Asp Tyr Ile Phe Pro Asp Thr Ser Ala Ile Ala 50 60

Thr Leu Val Ser Lys Gly Met Asn Ile Phe Arg Val Gln Phe Met Met 65 70 75 80

Glu Arg Leu Val Pro Asn Ser Met Thr Gly Ser Tyr Asp Asp Ala Tyr 85 90 95

Leu Asn Asn Leu Thr Thr Val Val Asn Ala Ile Ala Ala Gly Val 100 105 110

His Ala Ile Val Asp Pro His Asn Tyr Gly Arg Tyr Asn Asn Glu Ile 115 120 125

Ile Ser Ser Thr Ala Asp Phe Gln Thr Phe Trp Gln Asn Leu Ala Gly 130 140

Gln Phe Lys Asp Asn Asp Leu Val Ile Phe Asp Thr Asn Asn Glu Tyr 145 150 155 160

Asn Thr Met Asp Gln Thr Leu Val Leu Asp Leu Asn Gln Ala Ala Ile 165 170 175

Asp Gly Ile Arg Ala Ala Gly Ala Thr Ser Gln Tyr Ile Phe Ala Glu 180 185 190

Gly Asn Ser Trp Ser Gly Ala Trp Thr Trp Ala Asp Ile Asn Asp Asn Page 1

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Gly Ser Gln Cys Thr Gly Val Asp Ser Tyr Ser Gly Asp Thr Ile Ala 50 60

Trp His Thr Ser Trp Ser Trp Ser Gly Gly Ser Ser Ser Val Lys Ser 65 70 75 80

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Aspergillus aculeatus

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Gly Val Asn Trp Lys Leu Tyr Ser Gly Pro Asn Gly Asp Thr Thr Val 165 170 175

Tyr Ser Phe Val Ala Asp Ser Thr Thr Glu Ser Phe Ser Gly Asp Leu 180 185 190

Asn Asp Phe Phe Thr Tyr Leu Val Asp Asn Glu Gly Val Ser Asp Glu 195 200 205

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Val Gly Lys Gln Ile Gln Arg Gly Arg Lys Ile Ser Asp Ile Asn Ser 100 105

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Asp Phe Phe Asn Tyr Leu Glu Arg Asn His Gly Tyr Pro Ala Arg Glu 210 215 220

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Asn Phe Ile Thr Gly Lys Gly Lys Tyr Ala Met Leu Asp Pro His Asn Page 6

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165

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Gln Asn Leu Pro Gly Val Glu Gly Lys Asp Tyr Ile Trp Pro Asp Pro 50 60

Asn Thr Ile Asp Thr Leu Ile Ser Lys Gly Met Asn Ile Phe Arg Val 65 70 75 80

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Gln Lys Gly Ala Tyr Ala Val Val Asp Pro His Asn Tyr Gly Arg Tyr 115 120 125

Tyr Asn Ser Ile Ile Ser Ser Pro Ser Asp Phe Gln Thr Phe Trp Lys 130 140

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Asn Phe Ala Gln Ser Asn Gly Lys Leu Ile Arg Gly His Thr Leu Val 100 105 110

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Leu 320

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His Asp Ile Thr Tyr Ser Gly Ser Trp Thr Ser Thr Gly Asn Ser Asn 100 105 110

Ser Tyr Leu Ser Val Tyr Gly Trp Thr Thr Gly Pro Leu Val Glu Tyr 115 120 125 Page 13

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Gln Tyr Trp Ser Ile Arg Gln Thr Lys Arg Val Gly Gly Thr Val Thr 180 185 190

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Tyr Phe Tyr Ser Trp Trp Ser Asp Gly Gly Gln Val Gln Tyr Thr 50 55

Asn Leu Glu Gly Ser Arg Tyr Gln Val Arg Trp Arg Asn Thr Gly Asn 65 70 75 80

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Tyr Gly Gly Tyr Phe Asn Pro Gln Gly Asn Gly Tyr Leu Ala Val Tyr 100 105 110

Gly Trp Thr Arg Asn Pro Leu Val Glu Tyr Tyr Val Ile Glu Ser Tyr 115 120 125

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Thr Asp Gly Asp Gln Tyr Asp Ile Phe Val Ser Thr Arg Tyr Asn Gln 145 150 150 160

Pro Ser Ile Asp Gly Thr Arg Thr Phe Gln Gln Tyr Trp Ser Ile Arg 165 170 175

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Thr Ser Gln Trp Thr Pro Pro Pro Ala Gln Thr Ser Ser Asn Pro Pro 65 70 75 80

Pro Thr Gly Gly Gly Gly Asn Thr Leu His Glu Lys Phe Lys Ala 85 90 95

Arg Gly Lys Gln Tyr Phe Gly Thr Glu Ile Asp His Tyr His Leu Asn 100 105 110

Asn Asn Gln Leu Met Glu Ile Ala Arg Arg Glu Phe Gly Gln Ile Thr 115 120 125

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130 135 Phe Ser Phe Gly Asn Ala Asp Arg Val Val Asp Trp Ala Thr Ser Asn 145 150 155 160 Gly Lys Leu Ile Arg Gly His Thr Leu Leu Trp His Ser Gln Leu Pro 165 170 175 Gln Trp Val Gln Asn Ile Asn Asp Arg Asn Thr Leu Thr Gln Val Ile 180 185 190 Glu Asn His Val Arg Thr Val Met Thr Arg Tyr Lys Gly Lys Ile Phe 195 200 205 His Tyr Asp Val Val Asn Glu Ile Leu Asp Glu Asn Gly Gly Leu Arg 210 215 220 Asn Ser Val Phe Ser Arg Val Leu Gly Glu Asp Phe Val Gly Ile Ala 225 230 235 240 Phe Arg Ala Ala Arg Ala Ala Asp Pro Asp Ala Lys Leu Tyr Ile Asn 245 250 255 Asp Tyr Asn Leu Asp Ser Ala Asn Tyr Ala Lys Thr Arg Gly Met Ile 260 265 270 Asn Leu Val Asn Lys Trp Val Ser Gln Gly Val Pro Ile Asp Gly Ile 275 280 285 Thr Gln Ala His Leu Ala Gly Pro Gly Gly Trp Asn Pro Ala Ser 290 295 300 Gly Val Pro Ala Ala Leu Gln Ala Leu Ala Gly Ala Asn Val Lys Glu 305 310 315 320 Val Ala Ile Thr Glu Leu Asp Ile Gln Gly Ala Gly Ala Asn Asp Tyr 325 330 335 Val Thr Val Ala Asn Ala Cys Leu Asn Val Gln Lys Cys Val Gly Ile 340 345 350 Thr Val Trp Gly Val Ser Asp Arg Asp Thr Trp Arg Ser Asn Glu Asn 355 360 365 Leu Leu Tyr Asp Arg Asp Tyr Arg Pro Lys Ala Ala Tyr Asn Ala 370 375 380 Leu Met Asn Ala Leu

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Asp Ser Pro Gly Gln Arg Glu Arg Ala Gly Tyr Glu Asp Lys Tyr Ala 50 55 60

Gln Tyr Asp Gln Ile Met Trp Lys Ser Gly Glu Phe Gly Leu Thr Thr 65 70 75 80

Pro Thr Asn Gly Gln Lys Trp Leu Phe Thr Glu Pro Glu Arg Gly Val 85 90 95

Phe Asn Phe Thr Glu Gly Asp Ile Val Thr Asn Leu Ala Arg Lys His $100 \hspace{1cm} 105 \hspace{1cm} 110$

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Pro Trp Val Glu Ser Thr Glu Trp Thr Pro Glu Glu Leu Arg Gln Val 130 135 140

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Tyr Ala Trp Asp Val Val Asn Glu Ala Leu Asn Glu Asp Gly Thr Tyr 165 170 175

Arg Glu Ser Val Phe Tyr Lys Val Leu Gly Glu Asp Tyr Ile Lys Leu 180 185 190

Ala Phe Glu Thr Ala Ala Lys Val Asp Pro His Ala Lys Leu Tyr Tyr 195 200 205

Asn Asp Tyr Asn Leu Glu Ser Pro Ser Ala Lys Thr Glu Gly Ala Lys 210 215 220

Arg Ile Val Lys Met Leu Lys Asp Ala Gly Ile Arg Ile Asp Gly Val 225 230 235 240

Gly Leu Gln Ala His Leu Val Ala Glu Ser His Pro Thr Leu Asp Glu Page 17

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10429-000.ST25 250

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Leu Thr Glu Leu Asp Ile Arg Leu Ser Ile Pro Ala Asn Ala Thr Asn 275 280 285

Leu Ala Gln Gln Arg Glu Ala Tyr Lys Asn Val Val Gly Ala Cys Val 290 295 300

Gln Val Arg Gly Cys Ile Gly Val Glu Ile Trp Asp Phe Tyr Asp Pro 305 310 315 320

Phe Ser Trp Val Pro Ala Thr Phe Pro Gly Gln Gly Ala Pro Leu Leu 325 330 335

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Asn Gln Asn Tyr Arg Thr Ser Gly Asn Val Asn Tyr Ser Pro Thr Asp 50 55 60

Asn Gly Tyr Ser Val Ser Phe Ser Asn Ala Gly Asp Phe Val Val Gly 65 70 75 80

Lys Gly Trp Arg Thr Gly Ala Thr Arg Asn Ile Thr Phe Ser Gly Ser 85 90 95

Thr Gln His Thr Ser Gly Thr Val Leu Val Ser Val Tyr Gly Trp Thr 100 105 110 Page 18

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Ala Gly Ser Ala Gln Gly Glu Lys Leu Gly Thr Val Glu Ser Asp Gly 130 140

Gly Thr Tyr Glu Ile Trp Arg His Gln Gln Val Asn Gln Pro Ser Ile 145 150 155 160

Glu Gly Thr Ser Thr Phe Trp Gln Tyr Ile Ser Asn Arg Val Ser Gly 165 170 175

Gln Arg Pro Asn Gly Gly Thr Val Thr Leu Ala Asn His Phe Ala Ala 180 185 190

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Ser Gly 225

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Thr Tyr Glu Ile Ser Trp Gly Asp Gly Gly Asn Leu Val Gly Gly Lys 65 70 75 80

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Tyr Gln Pro Asn Gly Asn Ser Tyr Leu Ala Val Tyr Gly Trp Thr Arg 100 105 110

10429-000.ST25

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Pro Ser Ser Gly Ala Thr Asp Leu Gly Thr Val Glu Cys Asp Gly Ser 130 135 140

Ile Tyr Arg Leu Gly Lys Thr Thr Arg Val Asn Ala Pro Ser Ile Asp 145 150 155 160

Gly Thr Gln Thr Phe Asp Gln Tyr Trp Ser Val Arg Gln Asp Lys Arg 165 170 175

Thr Ser Gly Thr Val Gln Thr Gly Cys His Phe Asp Ala Trp Ala Arg 180 185 190

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Gln Gln Thr Gln Leu Cys Asp Gln Tyr Ala Thr Tyr Thr Gly Ser Val 20 25 30

Tyr Thr Ile Asn Asn Asn Leu Trp Gly Lys Asp Ala Gly Ser Gly Ser 35 40 45

Gln Cys Thr Thr Val Asn Ser Ala Ser Ser Ala Gly Thr Ser Trp Ser 50 60

Thr Lys Trp Asn Trp Ser Gly Gly Glu Asn Ser Val Lys Ser Tyr Ala 65 70 75 80

Asn Ser Gly Leu Ser Phe Asn Lys Lys Leu Val Ser Gln Ile Ser Arg 85 90 95

Ile Pro Thr Ala Ala Gln Trp Ser Tyr Asp Asn Thr Gly Ile Arg Ala 100 105 110

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Asp Val Ala Tyr Asp Leu Phe Thr Ala Ala Asp Ile Asn His Val Thr
115 120 125

Trp Ser Gly Asp Tyr Glu Leu Met Ile Trp Leu Ala Arg Tyr Gly Gly 130 140

Val Gln Pro Leu Gly Ser Lys Ile Ala Thr Ala Thr Val Glu Gly Gln 145 150 150 155

Thr Trp Glu Leu Trp Tyr Gly Val Asn Gly Ala Gln Lys Thr Tyr Ser 165 170 175

Phe Val Ala Pro Thr Pro Ile Thr Ser Phe Gln Gly Asp Val Asn Asp 180 185 190

Phe Phe Lys Tyr Leu Thr Gln Asn His Gly Phe Pro Ala Ser Ser Gln 195 200 205

Tyr Leu Ile Thr Leu Gln Phe Gly Thr Glu Pro Phe Thr Gly Gly Pro 210 215 220

Ala Thr Leu Thr Val Ser Asp Trp Ser Ala Ser Val Gln 225 230 235

<210> <211> 17 347

<212> PRT

T. reesei

<220>

<221> <222>

PEPTIDE (1)..(347)

<400>

Met Lys Ala Asn Val Ile Leu Cys Leu Leu Ala Pro Leu Val Ala Ala 1 10 15

Leu Pro Thr Glu Thr Ile His Leu Asp Pro Glu Leu Ala Ala Leu Arg 20 25 30

Ala Asn Leu Thr Glu Arg Thr Ala Asp Leu Trp Asp Arg Gln Ala Ser

Gln Ser Ile Asp Gln Leu Ile Lys Arg Lys Gly Lys Leu Tyr Phe Gly 50 60

Thr Ala Thr Asp Arg Gly Leu Leu Gln Arg Glu Lys Asn Ala Ala Ile 65 70 75 80

The Gin Ala Asp Leu Gly Gin Val Thr Pro Glu Asn Ser Met Lys Trp 85 90 95

Gln Ser Leu Glu Asn Asn Gln Gly Gln Leu Asn Trp Gly Asp Ala Asp 100 105 110 Tyr Leu Val Asn Phe Ala Gln Gln Asn Gly Lys Ser Ile Arg Gly His 115 120 125 Thr Leu Ile Trp His Ser Gln Leu Pro Ala Trp Val Asn Asn Ile Asn 130 140 Asn Ala Asp Thr Leu Arg Gln Val Ile Arg Thr His Val Ser Thr Val 145 150 160 Val Gly Arg Tyr Lys Gly Lys Ile Arg Ala Trp Asp Val Val Asn Glu 165 170 175 Ile Phe Asn Glu Asp Gly Thr Leu Arg Ser Ser Val Phe Ser Arg Leu 180 185 190 Leu Gly Glu Glu Phe Val Ser Ile Ala Phe Arg Ala Arg Asp Ala 195 200 205 Asp Pro Ser Ala Arg Leu Tyr Ile Asn Asp Tyr Asn Leu Asp Arg Ala 210 215 220 Asn Tyr Gly Lys Val Asn Gly Leu Lys Thr Tyr Val Ser Lys Trp Ile 225 230 235 240 Ser Gln Gly Val Pro Ile Asp Gly Ile Gly Ser Gln Ser His Leu Ser 245 250 255 Gly Gly Gly Ser Gly Thr Leu Gly Ala Leu Gln Gln Leu Ala Thr 260 265 270 Val Pro Val Thr Glu Leu Ala Ile Thr Glu Leu Asp Ile Gln Gly Ala 275 280 285 Pro Thr Thr Asp Tyr Thr Gln Val Val Gln Ala Cys Leu Ser Val Ser 290 295 300 Lys Cys Val Gly Ile Thr Val Trp Gly Ile Ser Asp Lys Asp Ser Trp 305 310 315 Arg Ala Ser Thr Asn Pro Leu Leu Phe Asp Ala Asn Phe Asn Pro Lys 325 330 335 Pro Ala Tyr Asn Ser Ile Val Gly Ile Leu Gln 340 345

<210> 18 <211> 419 <212> PRT

<213> T.reesei

10429-000.ST25

<220> **PEPTIDE** (1)..(419)<400> Met Asn Lys Pro Met Ser Ser Leu Leu Leu Ala Ala Thr Leu Leu Ala 1 10 15 Gly Gly Ser Ile Ala Gln Gln Thr Val Trp Gly Gln Cys Gly Gly Gln 20 25 30 Gly Trp Ser Gly Pro Thr Ser Cys Val Ala Gly Ser Ala Cys Ser Thr 35 40 45 Asn Pro Tyr Tyr Ala Gln Cys Ile Pro Gly Ala Thr Thr Met Ser 50 60Thr Thr Thr Lys Pro Thr Ser Val Ser Ala Ser Thr Thr Arg Ala Ser 65 70 75 80 Ala Thr Ser Ser Ala Thr Pro Pro Pro Ser Ser Gly Leu Thr Arg Phe 85 90 95 Ala Gly Val Asn Ile Ala Gly Phe Asp Phe Gly Cys Gly Thr Asp Gly 100 105 110 Thr Cys Val Thr Ser Lys Val Tyr Pro Pro Leu Lys Asn Tyr Ala Gly 115 120 125 Thr Asn Asn Tyr Pro Asp Gly Val Gly Gln Met Gln His Phe Val Asn 130 140 Asp Asp Lys Leu Thr Ile Phe Arg Leu Pro Val Gly Trp Gln Tyr Leu 145 150 155 160 Val Asn Asn Asn Leu Gly Gly Thr Leu Asp Ser Asn Asn Phe Gly Lys 165 170 175 Tyr Asp Gln Leu Val Gln Ala Cys Leu Ser Leu Gly Val Tyr Cys Ile 180 185 190 Val Asp Ile His Asn Tyr Ala Arg Trp Asn Gly Gly Ile Ile Gly Gln 195 200 205 Gly Gly Pro Thr Asn Asp Gln Phe Thr Ser Leu Trp Ser Gln Leu Ala 210 215 220 Gln Lys Tyr Ala Ser Gln Ser Lys Val Trp Phe Gly Ile Met Asn Glu 225 230 235 240

10429-000.ST25 Pro His Asp Val Asn Ile Asn Thr Trp Ala Thr Thr Val Gln Ala Val 245 250 255 Val Thr Ala Ile Arg Asn Ala Gly Ala Thr Ser Gln Phe Ile Ser Leu 260 265 270 Pro Gly Asn Asp Trp Gln Ser Ala Gly Ala Phe Ile Ser Asp Gly Ser 275 280 285 Ala Ala Leu Ser Gln Val Lys Asn Pro Asp Gly Ser Thr Pro Asn 290 295 300 Leu Ile Phe Asp Leu His Lys Tyr Leu Asp Ser Asp Asn Ser Gly Thr 305 310 315 320 His Ala Asp Cys Val Thr Asn Asn Val Asn Asp Ala Phe Ser Pro Val 325 330 335 Ala Thr Trp Leu Arg Gln Asn Asn Arg Gln Ala Ile Leu Thr Glu Thr 340 345 350 Gly Gly Gly Asn Thr Gln Ser Cys Ile Gln Tyr Leu Cys Gln Gln Phe 355 Gln Tyr Ile Asn Gln Asn Ser Asp Val Tyr Leu Gly Tyr Val Gly Trp 370 375 380 Gly Ala Gly Ser Phe Asp Ser Thr Tyr Ile Leu Thr Glu Thr Pro Thr 385 390 395 400 Gly Ser Gly Ser Ser Trp Thr Asp Thr Ser Leu Val Ser Ser Cys Ile 405 410 415

Ser Arg Lys

<210> 19 <211> 459 <212> PRT

<212> PRT <213> T.viride

<220>

<221> PEPTIDE <222> (1)..(459)

<400> 19

Met Ala Pro Ser Val Thr Leu Pro Leu Thr Thr Ala Ile Leu Ala Ile 1 10 15

Ala Arg Leu Val Ala Ala Gln Gln Pro Gly Thr Ser Thr Pro Glu Val 20 25 30

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His Pro Lys Leu Thr Thr Tyr Lys Cys Thr Lys Ser Gly Gly Cys Val
35 40 45 Ala Gln Asp Thr Ser Val Val Leu Asp Trp Asn Tyr Arg Trp Met His 50 60 Asp Ala Asn Tyr Asn Ser Cys Thr Val Asn Gly Gly Val Asn Thr Thr 65 70 75 80 Leu Cys Pro Asp Glu Ala Thr Cys Gly Lys Asn Cys Phe Ile Glu Gly Val Asp Tyr Ala Ala Ser Gly Val Thr Thr Ser Gly Ser Ser Leu Thr 100 105 110Met Asn Gln Tyr Met Pro Ser Ser Ser Gly Gly Tyr Ser Ser Val Ser 115 120 125 Pro Arg Leu Tyr Leu Leu Asp Ser Asp Gly Glu Tyr Val Met Leu Lys 130 135 140 Leu Asn Gly Gln Glu Leu Ser Phe Asp Val Asp Leu Ser Ala Leu Pro 145 150 155 160 Cys Gly Glu Asn Gly Ser Leu Tyr Leu Ser Gln Met Asp Glu Asn Gly
165 170 175 Gly Ala Asn Gln Tyr Asn Thr Ala Gly Ala Asn Tyr Gly Ser Gly Tyr 180 185 190 Cys Asp Ala Gln Cys Pro Val Gln Thr Trp Arg Asn Gly Thr Leu Asn 195 200 205 Thr Ser His Gln Gly Phe Cys Cys Asn Glu Met Asp Ile Leu Glu Gly 210 215 220 Asn Ser Arg Ala Asn Ala Leu Thr Pro His Ser Cys Thr Ala Thr Ala 225 230 235 Cys Asp Ser Ala Gly Cys Gly Phe Asn Pro Tyr Gly Ser Gly Tyr Lys 245 250 255Ser Tyr Tyr Gly Pro Gly Asp Thr Val Asp Thr Ser Lys Thr Phe Thr 260 265 270 Ile Ile Thr Gln Phe Asn Thr Asp Asn Gly Ser Pro Ser Gly Asn Leu 275 280 285 Val Gly Ile Thr Arg Lys Tyr Gln Gln Asn Gly Val Asp Ile Pro Ser

Ala Gln Pro Gly Gly Asp Thr Ile Ser Ser Cys Pro Ser Ala Ser Ala 305 310 315 320

Tyr Gly Gly Leu Ala Thr Met Gly Lys Ala Leu Ser Ser Gly Met Val 325 330 335

Leu Val Phe Ser Ile Trp Asn Asp Asn Ser Gln Tyr Met Asn Trp Leu 340 345 350

Asp Ser Gly Asn Ala Gly Pro Cys Ser Ser Thr Glu Gly Asn Pro Ser 355 360 365

Asn Ile Leu Ala Asn Asn Pro Asn Thr His Val Val Phe Ser Asn Ile 370 375 380

Arg Trp Gly Asp Ile Gly Ser Thr Thr Asn Ser Thr Ala Pro Pro 885 390 395 400

Pro Pro Ala Ser Ser Thr Thr Phe Ser Thr Thr Arg Arg Ser Ser Thr 405 410 415

Thr Ser Ser Ser Pro Ser Cys Thr Gln Thr His Trp Gly Gln Cys Gly 420 425 430

Gly Ile Gly Tyr Ser Gly Cys Lys Thr Cys Thr Ser Gly Thr Thr Cys 435 440 445

Gln Tyr Ser Asn Asp Tyr Tyr Ser Gln Cys Leu 450 455

<210> 20 <211> 232

<211> 232 <212> PRT

<213> T.reesei

<220>

<221> PEPTIDE

<222> (1)..(232)

<400> 20

Met Lys Phe Leu Gln Val Leu Pro Ala Leu Ile Pro Ala Ala Leu Ala 1 5 10 15

Gln Thr Ser Cys Asp Gln Trp Ala Thr Phe Thr Gly Asn Gly Tyr Thr 20 25 30

Val Ser Asn Asn Leu Trp Gly Ala Ser Ala Gly Ser Gly Phe Gly Cys 40 45

Val Thr Ala Val Ser Leu Ser Gly Gly Ala His Ala Asp Trp Gln Trp 50 60

Ser Gly Gly Gln Asn Asn Val Lys Ser Tyr Gln Asn Ser Gln Ile Ala 65 70 75 80

Ile Pro Gln Lys Arg Thr Val Asn Ser Ile Ser Ser Met Pro Thr Thr 85 90 95

Ala Ser Trp Ser Tyr Ser Gly Ser Asn Ile Arg Ala Asn Val Ala Tyr 100 105 110

Asp Leu Phe Thr Ala Ala Asn Pro Asn His Val Thr Tyr Ser Gly Asp 115 120 125

Tyr Glu Leu Met Ile Trp Leu Gly Lys Tyr Gly Asp Ile Gly Pro Ile 130 135 140

Gly Ser Ser Gln Gly Thr Val Asn Val Gly Gly Gln Ser Trp Thr Leu 145 150 155 160

Tyr Tyr Gly Tyr Asn Gly Ala Met Gln Val Tyr Ser Phe Val Ala Gln 165 170 175

Thr Asn Thr Thr Asn Tyr Ser Gly Asp Val Lys Asn Phe Phe Asn Tyr 180 185 190

Leu Arg Asp Asn Lys Gly Tyr Asn Ala Ala Gly Gln Tyr Val Leu Ser 195 200 205

Tyr Gln Phe Gly Thr Glu Pro Phe Thr Gly Ser Gly Thr Leu Asn Val 210 215

Ala Ser Trp Thr Ala Ser Ile Asn 225 230